# Letter to the Editor

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None.

# Letter by Menzaghi et al Regarding Article, "Plasma Levels of Fatty Acid–Binding Protein 4, Retinol-Binding Protein 4, High-Molecular-Weight Adiponectin, and Cardiovascular Mortality Among Men With Type 2 Diabetes: A 22-Year Prospective Study"

#### To the Editor:

We read with interest the article on the paradoxical association between high-molecular-weight (HMW) adiponectin and cardiovascular mortality in men with type 2 diabetes mellitus from the HPFS (Health Professional Follow-Up Study).<sup>1</sup> These data resemble those we have previously reported<sup>2</sup> in men with type 2 diabetes mellitus, comprising a large subset of the same HPSF and additional Italian individuals.<sup>2</sup>

To address whether such association is sustained by a causeeffect relationship, a genetic risk score (GRS) built on 19 single nucleotide polymorphisms previously associated with adiponectin levels (Mendelian randomization) was used.<sup>1</sup> Notably, GRS was not associated with cardiovascular mortality, thus speaking against a causal role of HMW adiponectin on it. This finding is in marked contrast with our recent report, showing a significant association between rs832354 (a single nucleotide polymorphism strongly associated with HMW adiponectin<sup>3</sup>) and cardiovascular mortality in type 2 diabetes mellitus patients of both sexes.<sup>4</sup> Of note, our association was entirely driven by males (ie, no association observed among females),<sup>5</sup> thus coincidentally reinforcing the belief of a sexual dimorphic effect of serum adiponectin on cardiovascular mortality<sup>2</sup> and further stressing the difference with Liu et al<sup>1</sup> report.

Here, we address some possible causes of such strident discrepancy.

First, differences in the clinical set may have played a role. In fact, although in HPFS only 50% patients had baseline cardiovascular disease, all our patients had established coronary artery disease.<sup>4</sup> That this difference may be important is suggested by the observation that when in HPFS participants with baseline cardiovascular events were excluded, the hazard ratio (95% confidence interval) of cardiovascular mortality of patients with high HMW adiponectin decreased from 2.07 (1.42–3.06) to 1.39 (0.54–3.61),<sup>1</sup> a 54% reduction in a log scale. It is, therefore, likely that in the absence of baseline cardiovascular mortality is attenuated, thus making difficult addressing causality.

Second, also differences in environmental and genetic background across studies have to be considered. In fact, HPFS patients were recruited from all over the United States, whereas ours were all recruited in a very small and homogeneous region of Central Southern Italy. In addition, whereas our participants were all Whites,<sup>4</sup> those of HPFS were not.<sup>1</sup> Because genetic background of serum adiponectin is heterogeneous across ethnicities,<sup>6</sup> such difference too may have contributed to discrepancy between the 2 studies. Third, statistical power is important to make reliable Mendelian randomization studies.<sup>7</sup> Unfortunately, in Liu et al<sup>1</sup> article, no power was reported for the association between GRS and cardiovascular mortality. Because GRS was barely associated with HMW adiponectin (P=0.02),<sup>1</sup> the risk of false-negative findings on its association with cardiovascular mortality has to be seriously considered.

Finally, it seems, also by the reference quoted (ie, 29, which reports a case–control study), that a logistic rather than a Cox regression was used when analyzing the association between GRS and cardiovascular mortality.<sup>1</sup> If this is the case, ignoring time to event could have altered the results and contributed to the discrepancy with our study.

In all, disentangling the intrinsic nature of the paradoxical association between adiponectin and cardiovascular mortality in type 2 diabetes mellitus is a difficult task; further collaborative studies, in extensively phenotyped patients of different ethnicities from wellpowered samples, are definitely needed to accomplish it.

# Acknowledgments

This communication was supported by grants from the Italian Ministry of Health grants, RC2014, RC2015, RC2016, and RF-2013-02356459, European Foundation for the Study of Diabetes/Pfizer grants and Società Italiana di Diabetologia-Fondazione Diabete Ricerca (C. Menzaghi).

### Disclosures

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<sup>(</sup>Arterioscler Thromb Vasc Biol. 2017;37:e55-e56. DOI: 10.1161/ATVBAHA.117.309308.) © 2017 American Heart Association, Inc.

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